

FIGURE 1





MRPSGTAGAA	LLALLAALCP	ASRALEEKKV	CQGTSNKLTQ	LGTFEDHFLS	-50
LQRMFNNCEV	VLGNLEITYV	QRNYDLSFLK	TIQEVAGYVL	IALNTVERIP	100
LENLQIIRGN	MYYENSYALA	VLSNYDANKT	GLKELPMRNL	QEILHGAVRF	150
SNNPALCNVE	SIQWRDIVSS	DFLSNMSMDF	QNHLGSCQKC	DPSCPNGSCW	200
GAGEENCQKL	TKIICAQQCS	GRCRGKSPSD	CCHNQCAAGC	TGPRESDCLV	250
CRKFRDEATC	KDTCPPLMLY	NPTTYQMDVN	PEGKYSFGAT	CVKKCPRNYV	300
VTDHGSCVRA	CGADSYEMEE	DGVRKCKKCE	GPCRKVCNGI	GIGEFKDSLS	350
INATNIKHFK	NCTSISGDLH	ILPVAFRGDS	${\tt FTHTPPLDPQ}$	ELDILKTVKE	400
ITGFLLIQAW	PENRTDLHAF	ENLEIIRGRT	KQHGQFSLAV	VSLNITSLGL	450
RSLKEISDGD	VIISGNKNLC	YANTINWKKL	FGTSGQKTKI	ISNRGENSCK	500
ATGQVCHALC	SPEGCWGPEP	RDCVSCRNVS	RGRECVDKCN	LLEGEPREFV	550
ENSECIQCHP	ECLPQAMNIT	CTGRGPDNCI	QCAHYIDGPH	CVKTCPAGVM	600
GENNTLVWKY	ADAGHVCHLC	HPNCTYGCTG	PGLEGCPTNG	PKIPSIATGM	650
VGALLLLLVV	ALGIGLFMRR	RHIVRKRTLR	RLLQERELVE	PLTPSGEAPN	700
QALLRILKET	EFKKIKVLGS	GAFGTVYKGL	WIPEGEKVKI	PVAIKELREA	750
TSPKANKEIL	DEAYVMASVD	NPHVCRLLGI	CLTSTVQLIT	QLMPFGCLLD	800
YVREHKDNIG	SQYLLNWCVQ	IAKGMNYLED	RRLVHRDLAA	RNVLVKTPQH	850
VKITDFGLAK	LLGAEEKEYH	AEGGKVPIKW	MALESILHRI	YTHQSDVWSY	900
GVTVWELMTF	GSKPYDGIPA	SEISSILEKG	ERLPQPPICT	IDVYMIMVKC	950
WMIDADSRPK	FRELIIEFSK	MARDPQRYLV	IQGDERMHLP	SPTDSNFYRA	1000
LMDEEDMDDV	VDADEYLIPQ	QGFFSSPSTS	RTPLLSSLSA	TSNNSTVACI	1050
DRNGLQSCPI	KEDSFLQRYS	SDPTGALTED	SIDDTFLPVP	EYINQSVPKR	1100
PAGSVQNPVY	HNQPLNPAPS	RDPHYQDPHS	TAVGNPEYLN	TVQPTCVNST	1150
FDSPAHWAQK	GSHQISLDNP	DYQQDFFPKE	AKPNGIFKGS	TAENAEYLRV	1200
APQSSEFIGA				,	1210
	LQRMFNNCEV LENLQIIRGN SNNPALCNVE GAGEENCQKL CRKFRDEATC VTDHGSCVRA INATNIKHFK ITGFLLIQAW RSLKEISDGD ATGQVCHALC ENSECIQCHP GENNTLVWKY VGALLLLLVV QALLRILKET TSPKANKEIL YVREHKDNIG VKITDFGLAK GVTVWELMTF WMIDADSRPK LMDEEDMDDV DRNGLQSCPI PAGSVQNPVY FDSPAHWAQK	LQRMFNNCEV VLGNLEITYV LENLQIIRGN MYYENSYALA SNNPALCNVE SIQWRDIVSS GAGEENCQKL TKIICAQQCS CRKFRDEATC KDTCPPLMLY VTDHGSCVRA CGADSYEMEE INATNIKHFK NCTSISGDLH ITGFLLIQAW PENRTDLHAF RSLKEISDGD VIISGNKNLC ATGQVCHALC SPEGCWGPEP ENSECIQCHP ECLPQAMNIT GENNTLVWKY ADAGHVCHLC VGALLLLLVV ALGIGLFMRR QALLRILKET EFKKIKVLGS TSPKANKEIL DEAYVMASVD YVREHKDNIG SQYLLNWCVQ VKITDFGLAK LLGAEEKEYH GVTVWELMTF GSKPYDGIPA WMIDADSRPK FRELIIEFSK LMDEEDMDDV VDADEYLIPQ DRNGLQSCPI KEDSFLQRYS PAGSVQNPVY HNQPLNPAPS	LQRMFNNCEV         VLGNLEITYV         QRNYDLSFLK           LENLQIIRGN         MYYENSYALA         VLSNYDANKT           SNNPALCNVE         SIQWRDIVSS         DFLSNMSMDF           GAGEENCQKL         TKIICAQQCS         GRCRGKSPSD           CRKFRDEATC         KDTCPPLMLY         NPTTYQMDVN           VTDHGSCVRA         CGADSYEMEE         DGVRKCKKCE           INATNIKHFK         NCTSISGDLH         ILPVAFRGDS           ITGFLLIQAW         PENRTDLHAF         ENLEIIRGRT           RSLKEISDGD         VIISGNKNLC         YANTINWKKL           ATGQVCHALC         SPEGCWGPEP         RDCVSCRNVS           ENSECIQCHP         ECLPQAMNIT         CTGRGPDNCI           GENNTLVWKY         ADAGHVCHLC         HPNCTYGCTG           VGALLLLLVV         ALGIGLFMRR         RHIVRKRTLR           QALLRILKET         EFKKIKVLGS         GAFGTVYKGL           TSPKANKEIL         DEAYVMASVD         NPHVCRLLGI           YVREHKDNIG         SQYLLNWCVQ         IAKGMNYLED           VKITDFGLAK         LLGAEEKEYH         AEGGKVPIKW           GVTVWELMTF         GSKPYDGIPA         SEISSILEKG           WMIDADSRPK         FRELIIEFSK         MARDPQRYLV           LMDEEDMDDV         VDADEYLIPQ         QGFFSSPSTS	LQRMFNNCEVVLGNLEITYVQRNYDLSFLKTIQEVAGYVLLENLQIIRGNMYYENSYALAVLSNYDANKTGLKELPMRNLSNNPALCNVESIQWRDIVSSDFLSNMSMDFQNHLGSCQKCGAGEENCQKLTKIICAQQCSGRCRGKSPSDCCHNQCAAGCCRKFRDEATCKDTCPPLMLYNPTTYQMDVNPEGKYSFGATVTDHGSCVRACGADSYEMEEDGVRKCKKCEGPCRKVCNGIINATNIKHFKNCTSISGDLHILPVAFRGDSFTHTPPLDPQITGFLLIQAWPENRTDLHAFENLEIIRGRTKQHGQFSLAVRSLKEISDGDVIISGNKNLCYANTINWKKLFGTSGQKTKIATGQVCHALCSPEGCWGPEPRDCVSCRNVSRGRECVDKCNENSECIQCHPECLPQAMNITCTGRGPDNCIQCAHYIDGPHGENNTLVWKYADAGHVCHLCHPNCTYGCTGPGLEGCPTNGVGALLILLVVALGIGLFMRRRHIVRKRTLRRLLQERELVEQALLRILKETEFKKIKVLGSGAFGTVYKGLWIPEGEKVKITSPKANKEILDEAYVMASVDNPHVCRLLGICLTSTVQLITYVREHKDNIGSQYLLNWCVQIAKGMNYLEDRRLVHRDLAAVKITDFGLAKLLGAEEKEYHAEGGKVPIKWMALESILHRIGVTVWELMTFGSKPYDGIPASEISSILEKGERLPQPPICTWMIDADSRPKFRELIIEFSKMARDPQRYLVIQGDERMHLPLMDEEDMDDVVDADEYLIPQQGFFSSPSTSRTPLLSSLSADRNGLQSCPIKEDSFLQRYSSDPTGALTEDSIDDTFLPVPPAGSVQNPVYHNQPLNPAPSRDPHYQDPHSTAVGNPEYLNFDSPAHWAQKGSHQISLDNPDYQQDFFPKEAKPNGIFKGS	MRPSGTAGAA LLALLAALCP ASRALEEKKV CQGTSNKLTQ LGTFEDHFLS LQRMFNNCEV VLGNLEITYV QRNYDLSFLK TIQEVAGYVL 1ALNTVERIP LENLQIIRGN MYYENSYALA VLSNYDANKT GLKELPMRNL QEILHGAVRF SNNPALCNVE SIQWRDIVSS DFLSNMSMDF QNHLGSCQKC DPSCPNGSCW GAGEENCQKL TKIICAQQCS GRCRGKSPSD CCHNQCAAGC TGPRESDCLV CRKFRDEATC KDTCPPLMLY NPTTYQMDVN PEGKYSFGAT CVKKCPRNYV VTDHGSCVRA CGADSYEMEE DGVRKCKKCE GPCRKVCNGI GIGEFKDSLS INATNIKHFK NCTSISGDLH ILPVAFRGDS FTHTPPLDPQ ELDILKTVKE ITGFLLIQAW PENRTDLHAF ENLEIIRGRT KQHGQFSLAV VSLNITSLGL RSLKEISDGD VIISGNKNLC YANTINWKKL FGTSGQKTKI ISNRGENSCK ATGQVCHALC SPEGCWGPEP RDCVSCRNVS RGRECVDKCN LLEGEPREFV ENSECIQCHP ECLPQAMNIT CTGRGPDNCI QCAHYIDGPH CVKTCPAGVM GENNTLVWKY ADAGHVCHLC HPNCTYGCTG PGLEGCPTNG PKIPSIATGM VGALLLLLVV ALGIGLFMRR RHIVKRRTLR RLLQERELVE PLTPSGEAPN QALLRILKET EFKKIKVLGS GAFGTVYKGL WIPEGEKVKI PVAIKELREA TSPKANKEIL DEAYVMASVD NPHVCRLLGI CLTSTVQLIT QLMPFGCLLD VVREHKDNIG SQYLLNWCVQ IAKGMNYLED RRLVHRDLAA RNVLVKTPQH VKITDFGLAK LLGAEEKEYH AEGGKVPIKW MALESILHRI YTHQSDVWSY GVTVWELMTF GSKPYDGIPA SEISSILEKG ERLPQPPICT IDVYMIMVKC WMIDADSRPK FRELIIEFSK MARDPQRYLV IQGDERMHLP SPTDSNFYRA LMDEEDMDDV VDADEYLIPQ QGFFSSPSTS RTPLLSSLSA TSNNSTVACI DRNGLQSCPI KEDSFLQRYS SDPTGALTED SIDDTFLPVP EYINQSVPKR PAGSVQNPVY HNQPLNPAPS RDPHYQDPHS TAVGNPEYLN TVQPTCVNST PAGSSEFIGA



	<b>ATG</b> CGACCCT	CCGGGACGGC	CGGGGCAGCG	CTCCTGGCGC	TGCTGGCTGC	50
	GCTCTGCCCG	GCGAGTCGGG	CTCTGGAGGA	AAAGAAAGTT	TGCCAAGGCA	100
	CGAGTAACAA	GCTCACGCAG	TTGGGCACTT	TTGAAGATCA	TTTTCTCAGC	150
5	CTCCAGAGGA	TGTTCAATAA	CTGTGAGGTG	GTCCTTGGGA	ATTTGGAAAT	200
	TACCTATGTG	CAGAGGAATT	ATGATCTTTC	CTTCTTAAAG	ACCATCCAGG	250
	AGGTGGCTGG	TTATGTCCTC	ATTGCCCTCA	ACACAGTGGA	GCGAATTCCT	300
	TTGGAAAACC	TGCAGATCAT	CAGAGGAAAT	ATGTACTACG	AAAATTCCTA	350
	TGCCTTAGCA	GTCTTATCTA	ACTATGATGC	АААТААААСС	GGACTGAAGG	400
10	AGCTGCCCAT	GAGAAATTTA	CAGGAAATCC	TGCATGGCGC	CGTGCGGTTC	450
	AGCAACAACC	CTGCCCTGTG	CAACGTGGAG	AGCATCCAGT	GGCGGGACAT	500
	AGTCAGCAGT	GACTTTCTCA	GCAACATGTC	GATGGACTTC	CAGAACCACC	550
	TGGGCAGCTG	CCAAAAGTGT	GATCCAAGCT	GTCCCAATGG	GAGCTGCTGG	600
	GGTGCAGGAG	AGGAGAACTG	CCAGAAACTG	ACCAAAATCA	TCTGTGCCCA	650
15	GCAGTGCTCC	GGGCGCTGCC	GTGGCAAGTC	CCCCAGTGAC	TGCTGCCACA	700
	ACCAGTGTGC	TGCAGGCTGC	ACAGGCCCCC	GGGAGAGCGA	CTGCCTGGTC	750
	TGCCGCAAAT	TCCGAGACGA	AGCCACGTGC	AAGGACACCT	GCCCCCACT	800
	CATGCTCTAC	AACCCCACCA	CGTACCAGAT	GGATGTGAAC	CCCGAGGGCA	850
	AATACAGCTT	TGGTGCCACC	TGCGTGAAGA	AGTGTCCCCG	TAATTATGTG	900
20	GTGACAGATC	ACGGCTCGTG	CGTCCGAGCC	TGTGGGGCCG	ACAGCTATGA	950
	GATGGAGGAA	GACGGCGTCC	GCAAGTGTAA	GAAGTGCGAA	GGGCCTTGCC	1000
	GCAAAGTGTG	TAACGGAATA	GGTATTGGTG	AATTTAAAGA	CTCACTCTCC	1050
•	ATAAATGCTA	CGAATATTAA	ACACTTCAAA	AACTGCACCT	CCATCAGTGG	1100
	CGATCTCCAC	ATCCTGCCGG	TGGCATTTAG	GGGTGACTCC	TTCACACATA	1150
25	CTCCTCCTCT	GGATCCACAG	GAACTGGATA	TTCTGAAAAC	CGTAAAGGAA	1200
	ATCACAGGGT	TTTTGCTGAT	TCAGGCTTGG	CCTGAAAACA	GGACGGACCT	1250
	CCATGCCTTT	GAGAACCTAG	AAATCATACG	CGGCAGGACC	AAGCAACATG	1300
	GTCAGTTTTC	TCTTGCAGTC	GTCAGCCTGA	ACATAACATC	CTTGGGATTA	1350
	CGCTCCCTCA	AGGAGATAAG	TGATGGAGAT	GTGATAATTT	CAGGAAACAA	1400



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	AAATTTGTGC TATGCAAAT	А СААТАААСТО	GAAAAAACTG	TTTGGGACCT	1450
	CCGGTCAGAA AACCAAAAT	T ATAAGCAACA	GAGGTGAAAA	CAGCTGCAAG	1500
	GCCACAGGCC AGGTCTGCC	A TGCCTTGTGC	TCCCCGAGG	GCTGCTGGGG	1550
5	CCCGGAGCCC AGGGACTGC	G TCTCTTGCCG	GAATGTCAGC	CGAGGCAGGG	1600
	AATGCGTGGA CAAGTGCAAG	G CTTCTGGAGG	GTGAGCCAAG	GGAGTTTGTG	1650
•	GAGAACTCTG AGTGCATACA	A GTGCCACCCA	GAGTGCCTGC	CTCAGGCCAT	1700
	GAACATCACC TGCACAGGA	C GGGGACCAGA	CAACTGTATC	CAGTGTGCCC	1750
	ACTACATTGA CGGCCCCCAC	C TGCGTCAAGA	CCTGCCCGGC	AGGAGTCATG	1800
10	GGAGAAAACA ACACCCTGGT	CTGGAAGTAC	GCAGACGCCG	GCCATGTGTG	1850
	CCACCTGTGC CATCCAAACT	GCACCTACGG	ATGCACTGGG	CCAGGTCTTG	1900
	AAGGCTGTCC AACGAATGGC	CCTAAGATCC	CGTCCATCGC	CACTGGGATG	1950
	GTGGGGGCCC TCCTCTTGCT	GCTGGTGGTG	GCCCTGGGGA	TCGGCCTCTT	2000
	CATGCGAAGG CGCCACATCG	TTCGGAAGCG	CACGCTGCGG	AGGCTGCTGC	2050
15	AGGAGAGGGA GCTTGTGGAG	CCTCTTACAC	CCAGTGGAGA	AGCTCCCAAC	2100
	CAAGCTCTCT TGAGGATCTT	GAAGGAAACT	GAATTCAAAA	AGATCAAAGT	2150
	GCTGGGCTCC GGTGCGTTCG	GCACGGTGTA	TAAGGGACTC	TGGATCCCAG	2200
	AAGGTGAGAA AGTTAAAATT	CCCGTCGCTA	TCAAGGAATT	AAGAGAAGCA	2250
	ACATCTCCGA AAGCCAACAA	GGAAATCCTC	GATGAAGCCT	ACGTGATGGC	2300
20	CAGCGTGGAC AACCCCCACG	TGTGCCGCCT	GCTGGGCATC	TGCCTCACCT	2350
	CCACCGTGCA ACTCATCACG	CAGCTCATGC	CCTTCGGCTG	CCTCCTGGAC	2400
	TATGTCCGGG AACACAAAGA	CAATATTGGC	TCCCAGTACC	TGCTCAACTG	2450
	GTGTGTGCAG ATCGCAAAGG	GCATGAACTA	CTTGGAGGAC	CGTCGCTTGG	2500
	TGCACCGCGA CCTGGCAGCC	AGGAACGTAC	TGGTGAAAAC	ACCGCAGCAT	2550
25	GTCAAGATCA CAGATTTTGG	GCTGGCCAAA	CTGCTGGGTG	CGGAAGAGAA	2600
	AGAATACCAT GCAGAAGGAG	GCAAAGTGCC	TATCAAGTGG	ATGGCATTGG	2650
	AATCAATTTT ACACAGAATC	TATACCCACC	AGAGTGATGT (	CTGGAGCTAC	2700
	GGGGTGACCG TTTGGGAGTT	GATGACCTTT	GGATCCAAGC	CATATGACGG	2750
	AATCCCTGCC AGCGAGATCT	CCTCCATCCT	GGAGAAAGGA (	GAACGCCTCC	2800
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DEMARK	CTCAGCCACC	CATATGTACC	ATCGATGTCT	ACATGATCAT	GGTCAAGTGC	2850
UE	TGGATGATAG	ACGCAGATAG	TCGCCCAAAG	TTCCGTGAGT	TGATCATCGA	2900
	ATTCTCCAAA	ATGGCCCGAG	ACCCCAGCG	${\tt CTACCTTGTC}$	ATTCAGGGGG	2950
. 5	ATGAAAGAAT	GCATTTGCCA	AGTCCTACAG	ACTCCAACTT	CTACCGTGCC	3000
	CTGATGGATG	AAGAAGACAT	GGACGACGTG	GTGGATGCCG	ACGAGTACCT	3050
	CATCCCACAG	CAGGGCTTCT	TCAGCAGCCC	CTCCACGTCA	CGGACTCCCC	3100
	TCCTGAGCTC	TCTGAGTGCA	ACCAGCAACA	ATTCCACCGT	GGCTTGCATT	3150
	GATAGAAATG	GGCTGCAAAG	CTGTCCCATC	AAGGAAGACA	GCTTCTTGCA	3200
10	GCGATACAGC	TCAGACCCCA	CAGGCGCCTT	GACTGAGGAC	AGCATAGACG	3250
	ACACCTTCCT	CCCAGTGCCT	GAATACATAA	ACCAGTCCGT	TCCCAAAAGG	3300
	CCCGCTGGCT	CTGTGCAGAA	TCCTGTCTAT	CACAATCAGC	CTCTGAACCC	3350
		AGAGACCCAC				3400
	GCAACCCCGA	GTATCTCAAC	ACTGTCCAGC	CCACCTGTGT	CAACAGCACA	3450
15	TTCGACAGCC	CTGCCCACTG	GGCCCAGAAA	GGCAGCCACC	AAATTAGCCT	3500
	GGACAACCCT	GACTACCAGC	AGGACTTCTT	TCCCAAGGAA	GCCAAGCCAA	3550
	ATGGCATCTT	TAAGGGCTCC	ACAGCTGAAA	ATGCAGAATA	CCTAAGGGTC	3600
	GCGCCACAAA	GCAGTGAATT	TATTGGAGCA	TGA		3630

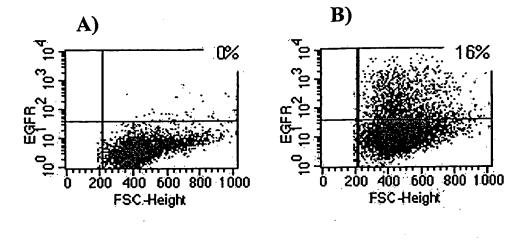


## First PCR Reaction

EG	FR1 + EGFR3 primer	
a.)		<b>E</b> GFR cDNA
	PCR product a encoding aa 1 to 24 fused to aa 313 to 319	
	EGFR2 + EGFR2220R primer	
b.)		<b>≡</b> EGFR cDNA
	PCR product b encoding aa 18 to 24 fus	sed to aa 313 to 678
Sec	cond PCR Reaction	
	PCR product a + b	
a.)	overlap extension	
	PCR product c encoding aa 1 to 24 fused to aa 3 with a stop codon in position 679	
	EGFR1 + EGFR2220R primer	
<b>D.</b> )	amplification of	

FIGURE 4





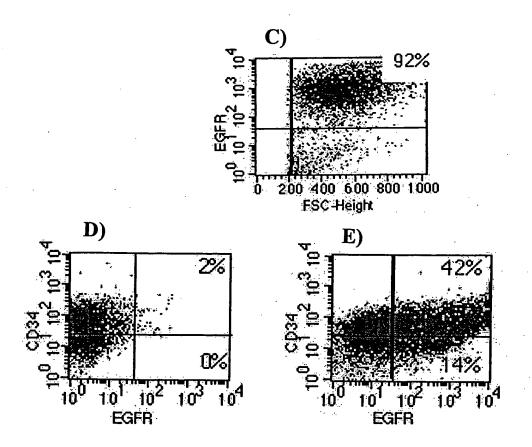


FIGURE 5